

Figure 1

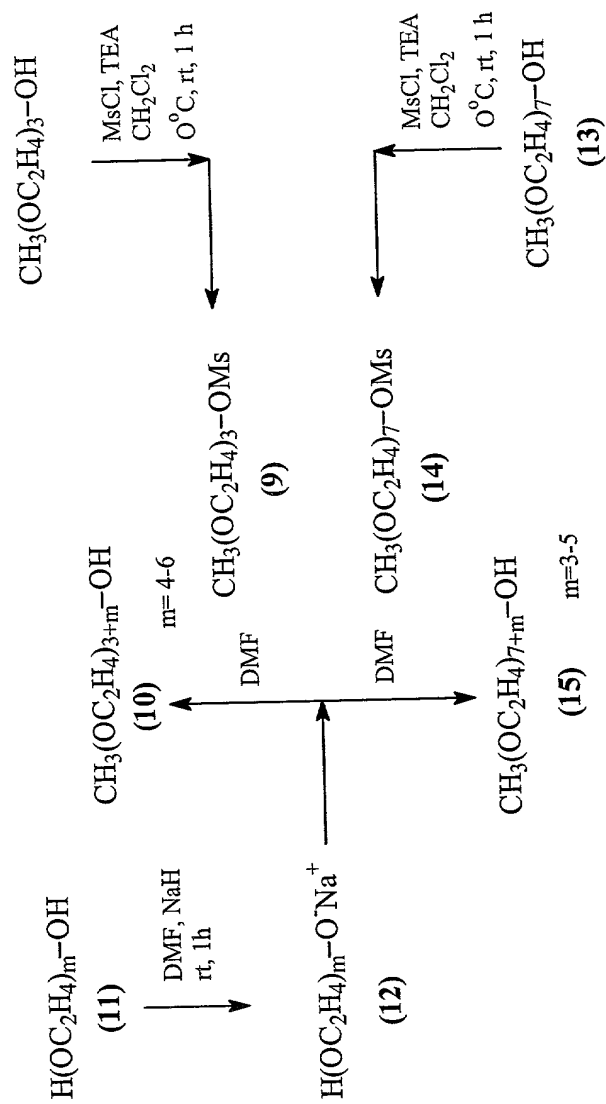


Figure 2

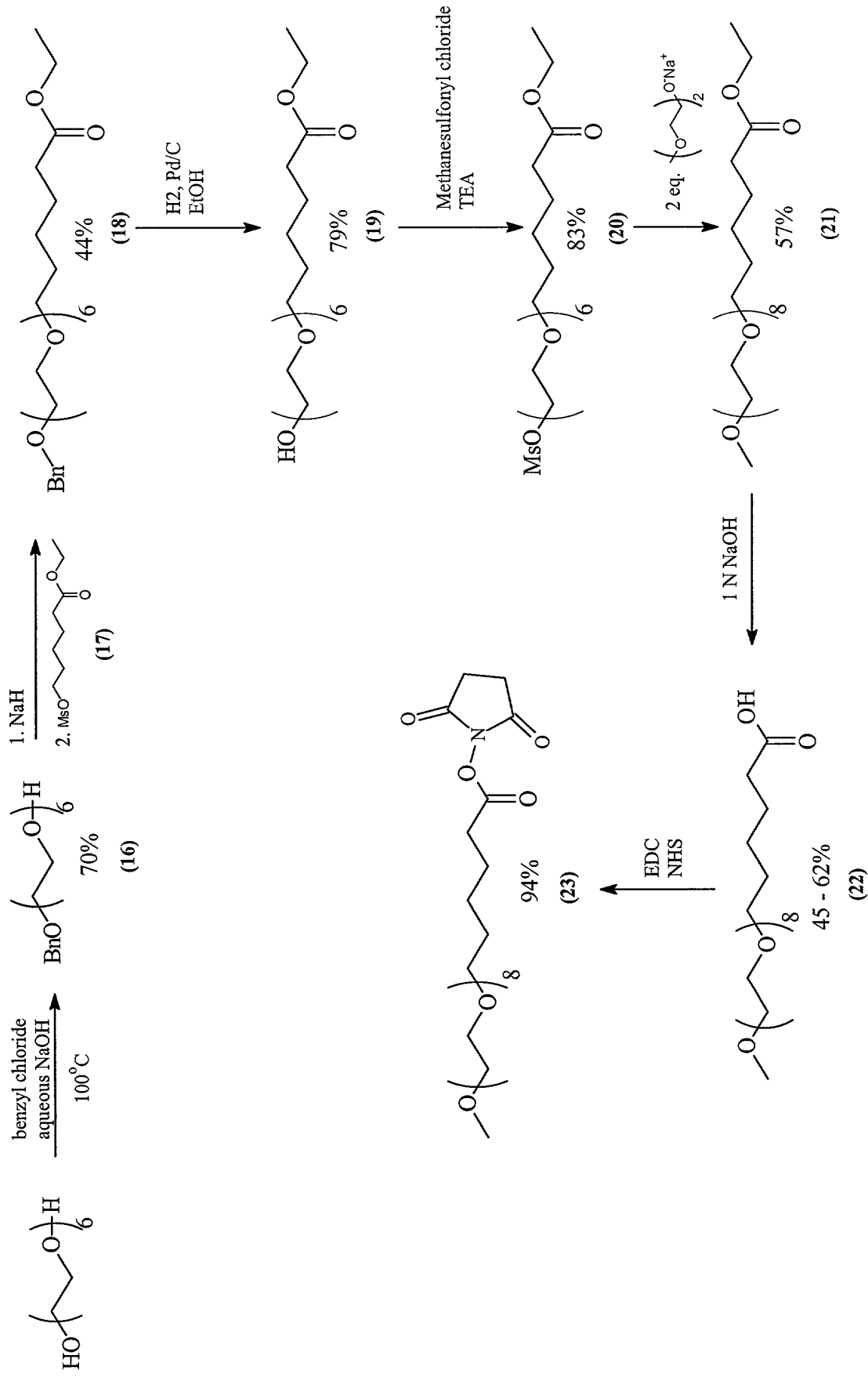


Figure 3

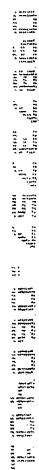


Figure 4

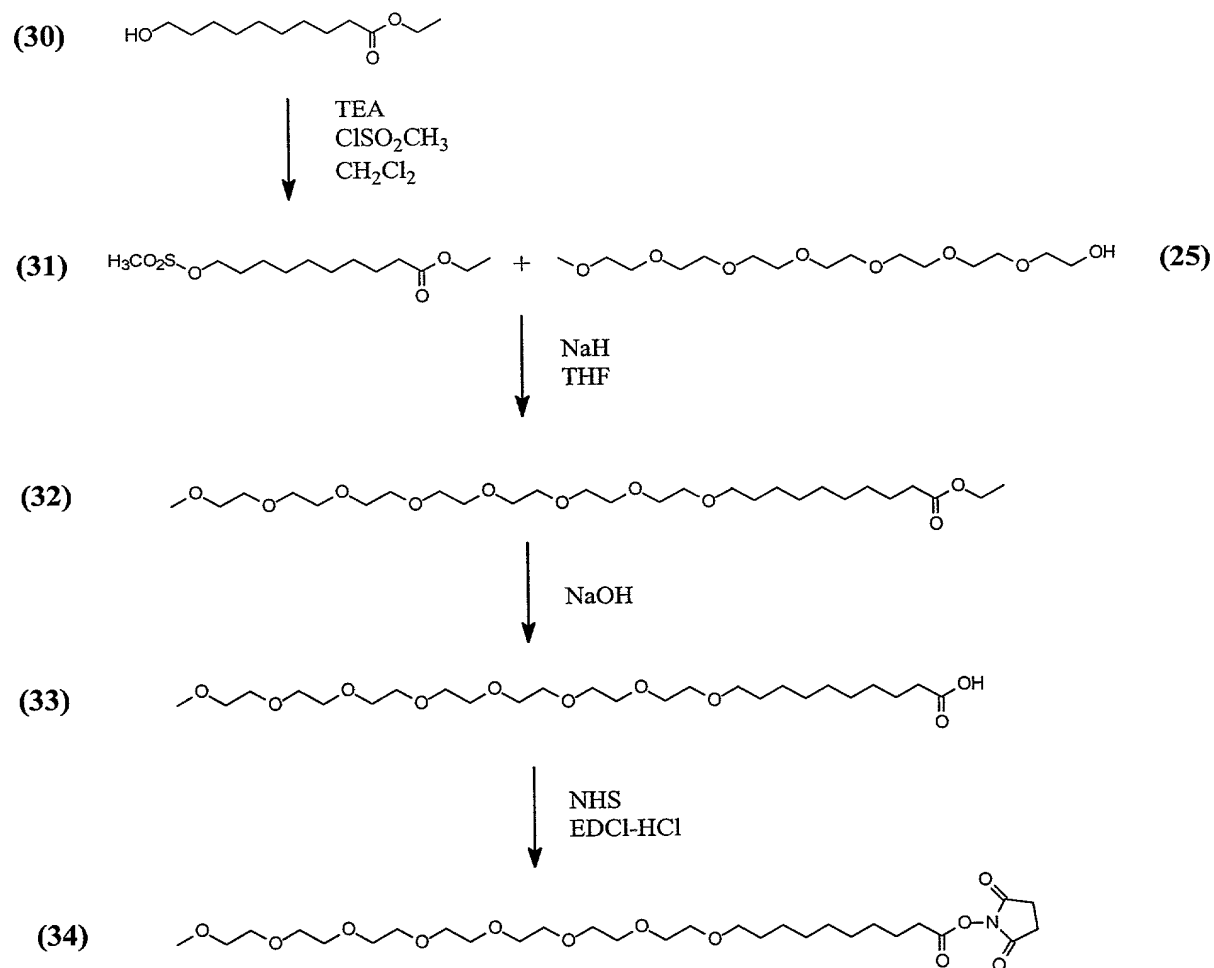


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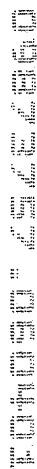
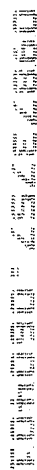


Figure 6



1000

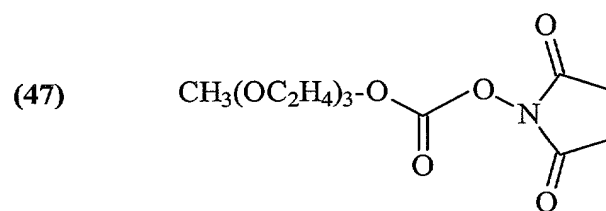
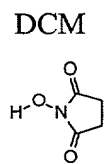
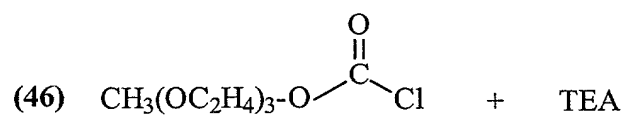
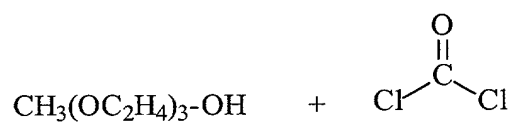


Figure 8

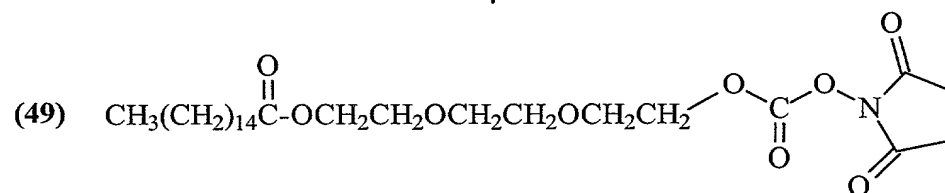
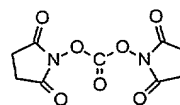
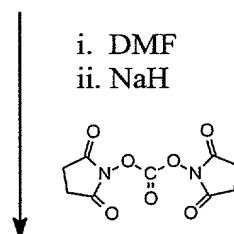
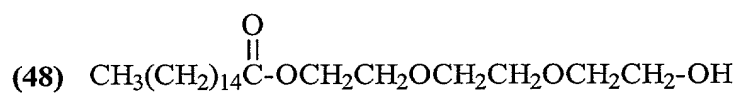
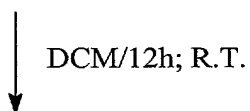
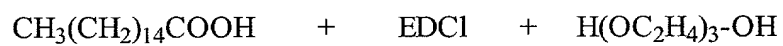
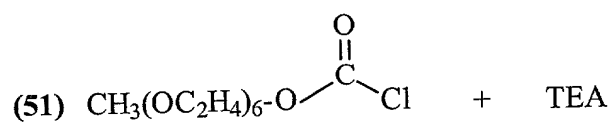
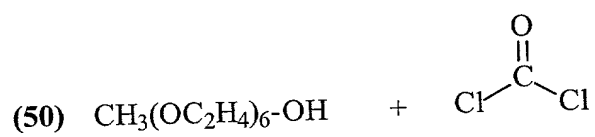


Figure 9



DCM

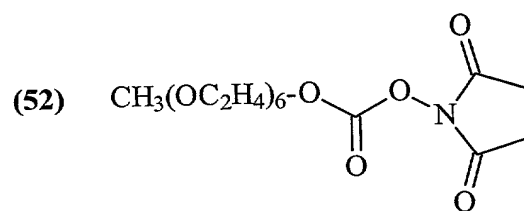
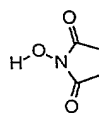


Figure 10

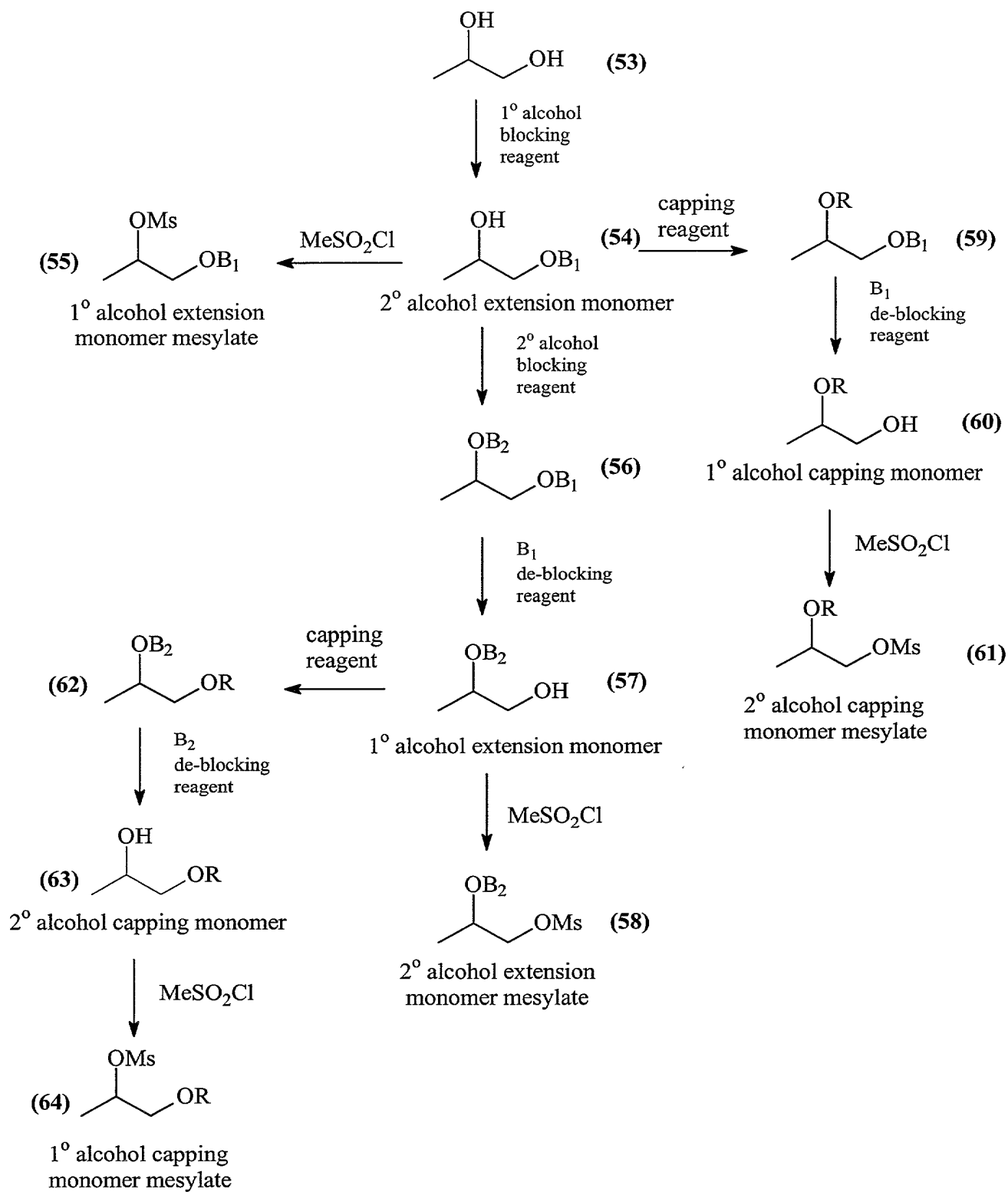


Figure 11

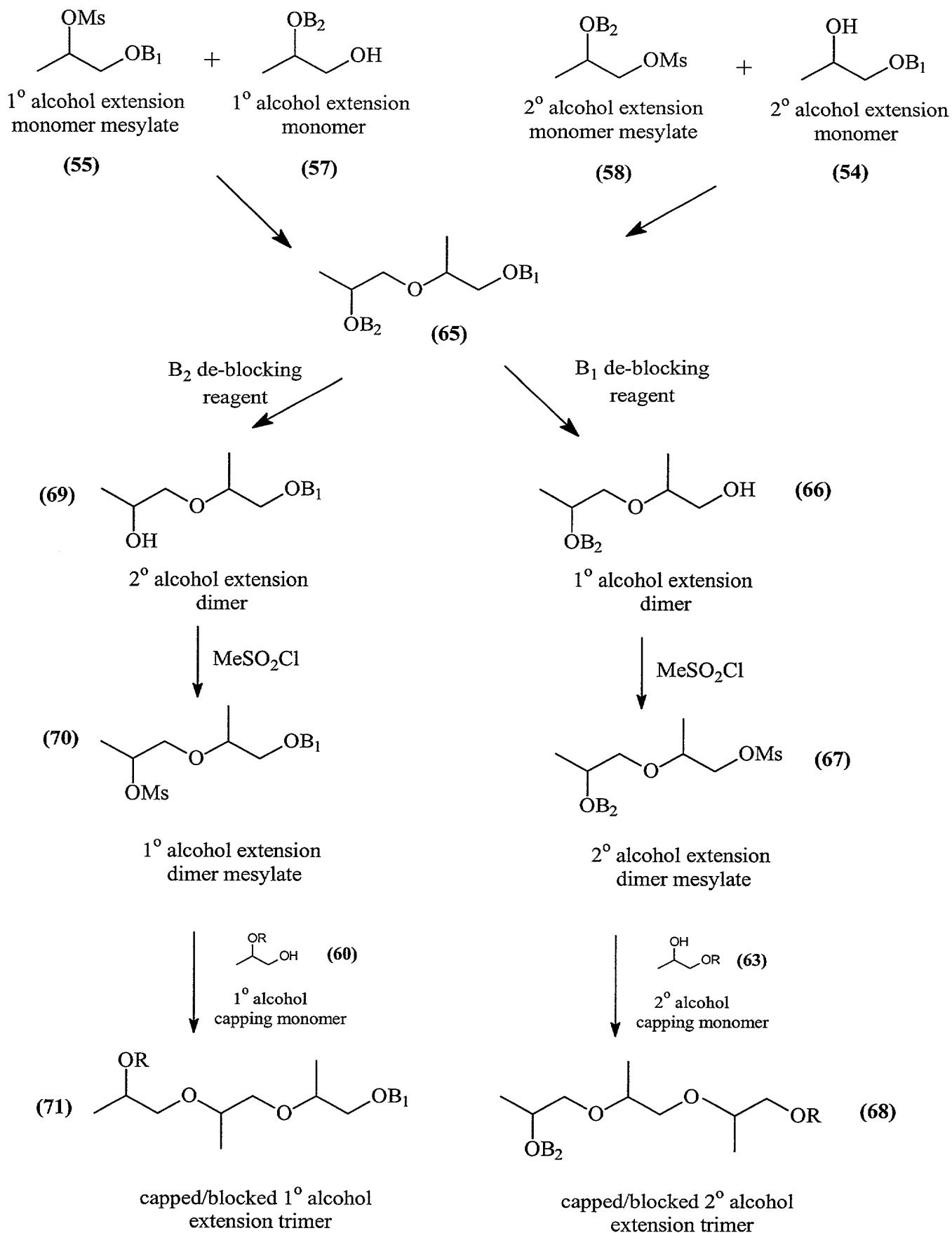


Figure 12

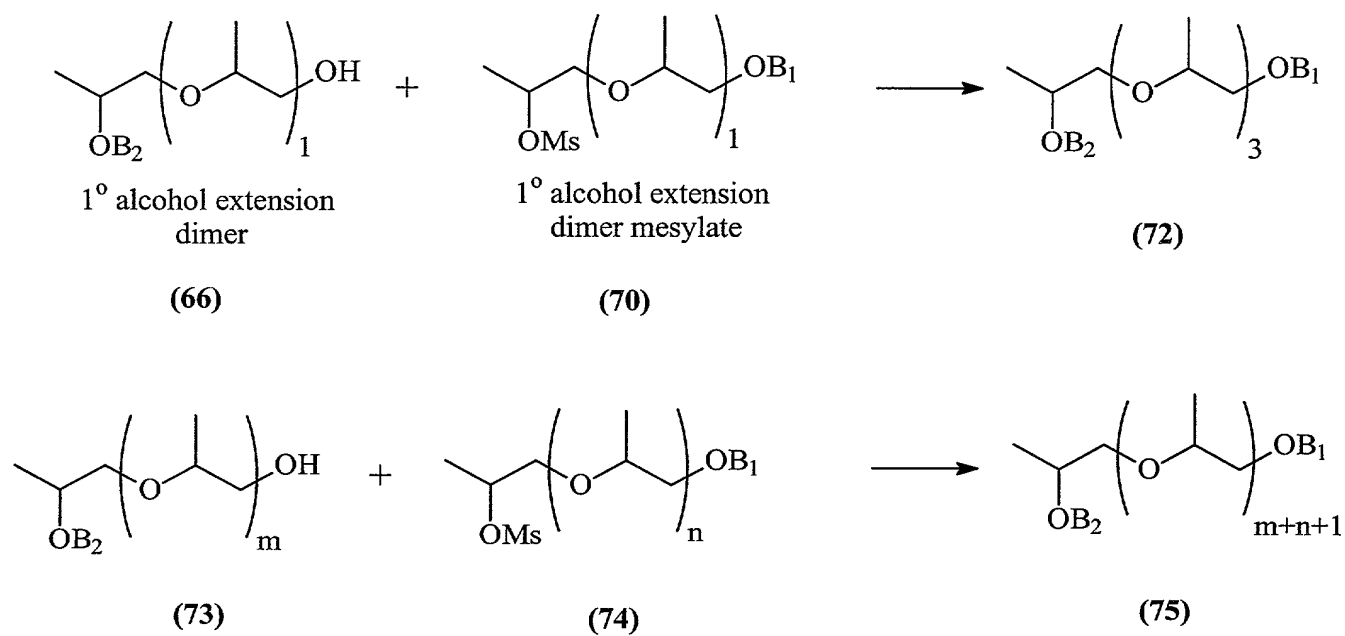


Figure 13

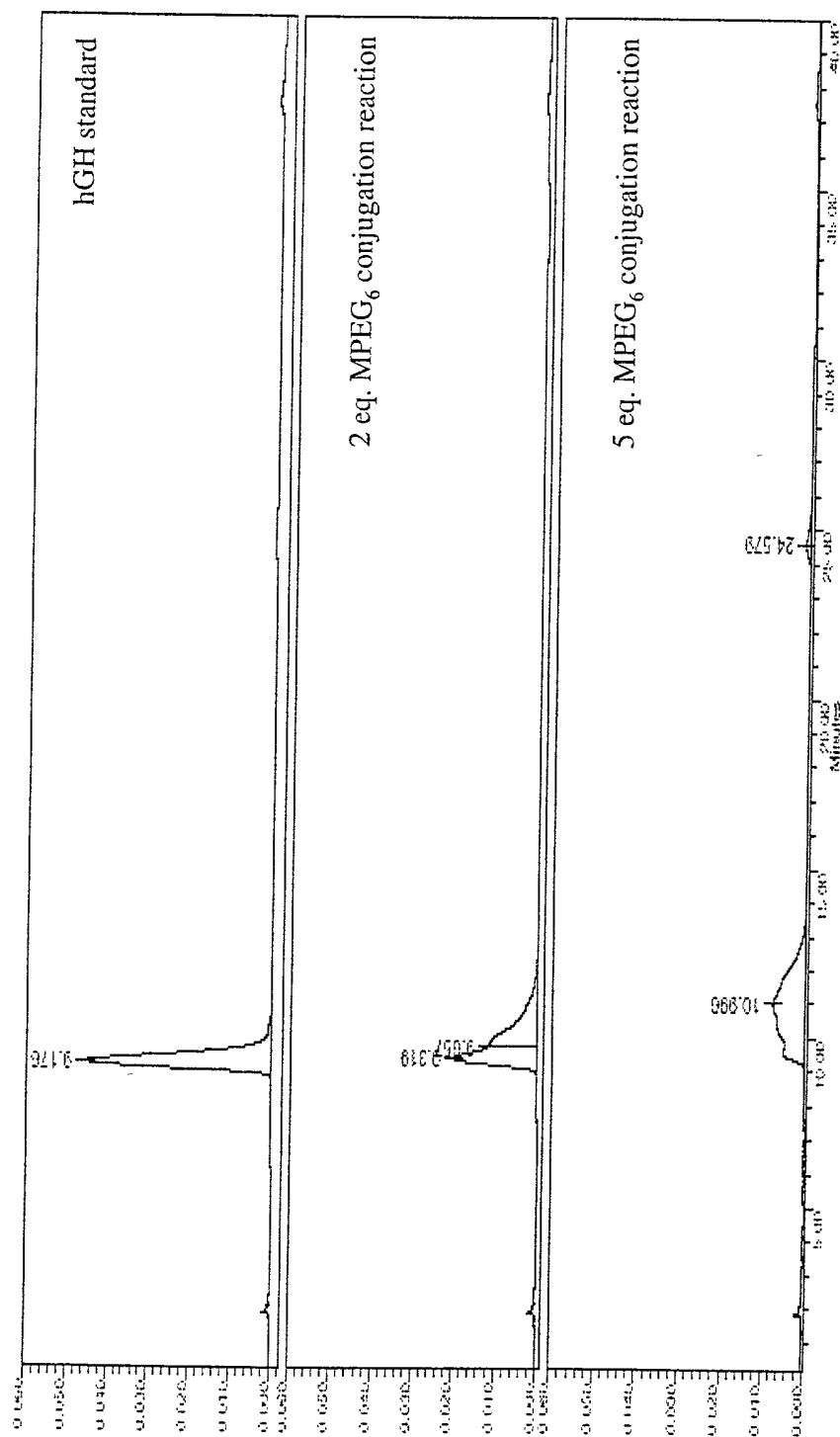


Figure 14

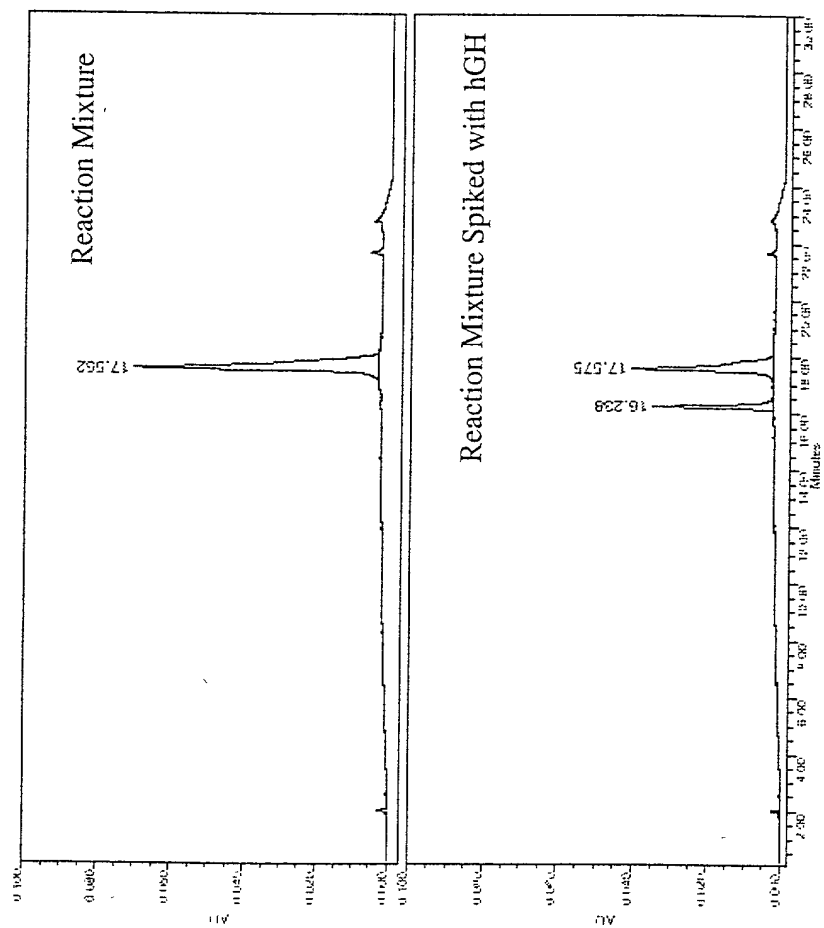


Figure 15

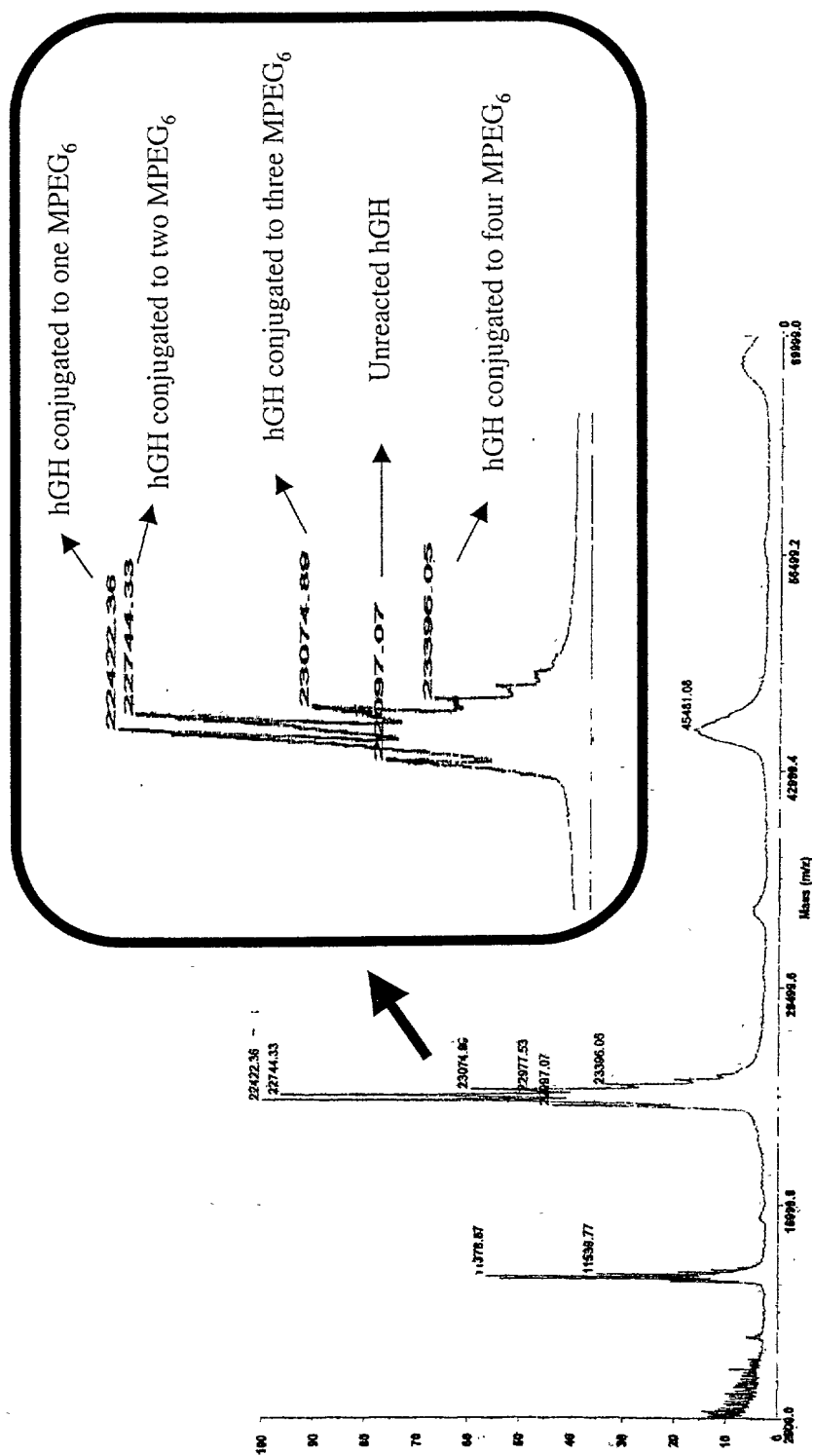
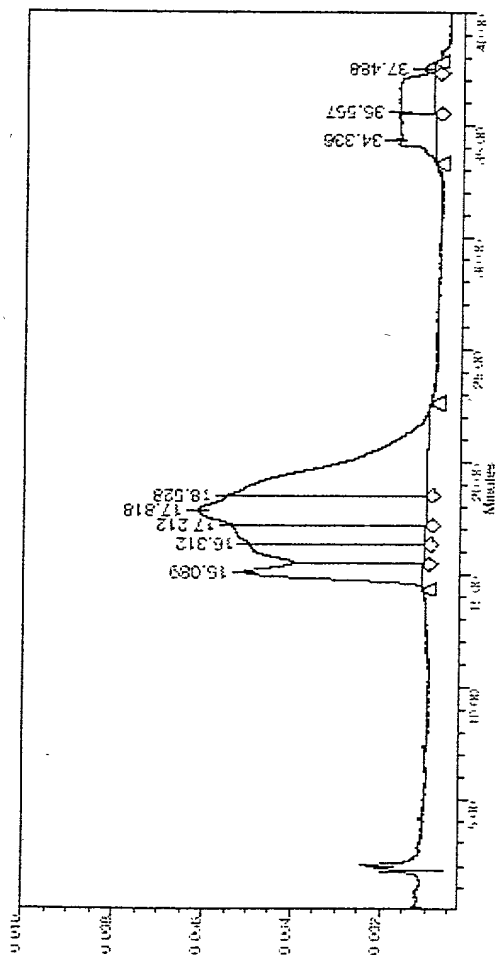


Figure 16



HPLC gradient: 50% to 70% acetonitrile in 30 minutes

Fraction	Retention Time (min.)	Area
150-AO-109A	15.089	155714
150-AO-109B	16.312	179786
150-AO-109C	17.212	221177
150-AO-109D	17.818	379932
150-AO-109E	18.528	400256
150-AO-109F	34.336	74735
150-AO-109G	35.557	76637

Figure 17

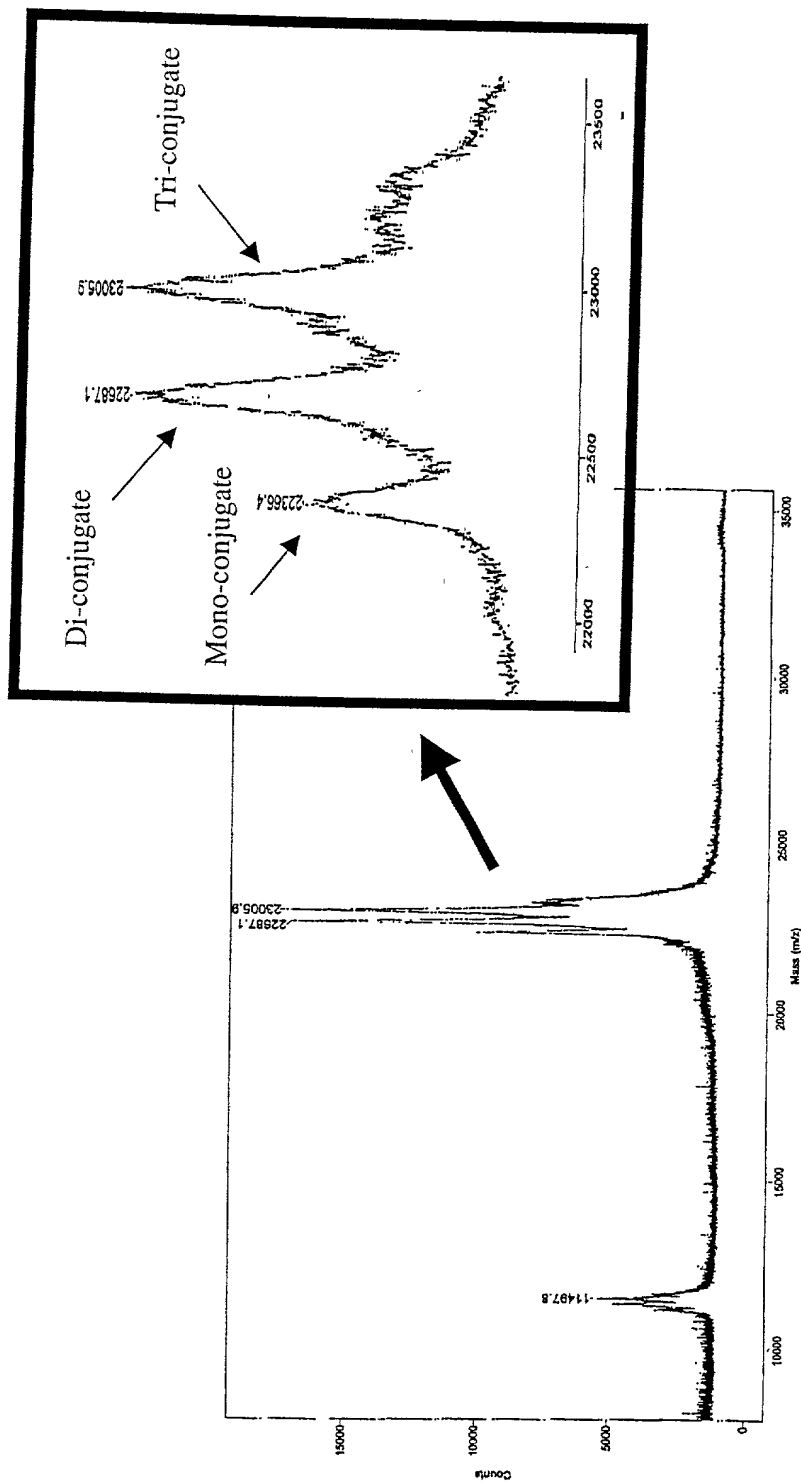


Figure 18

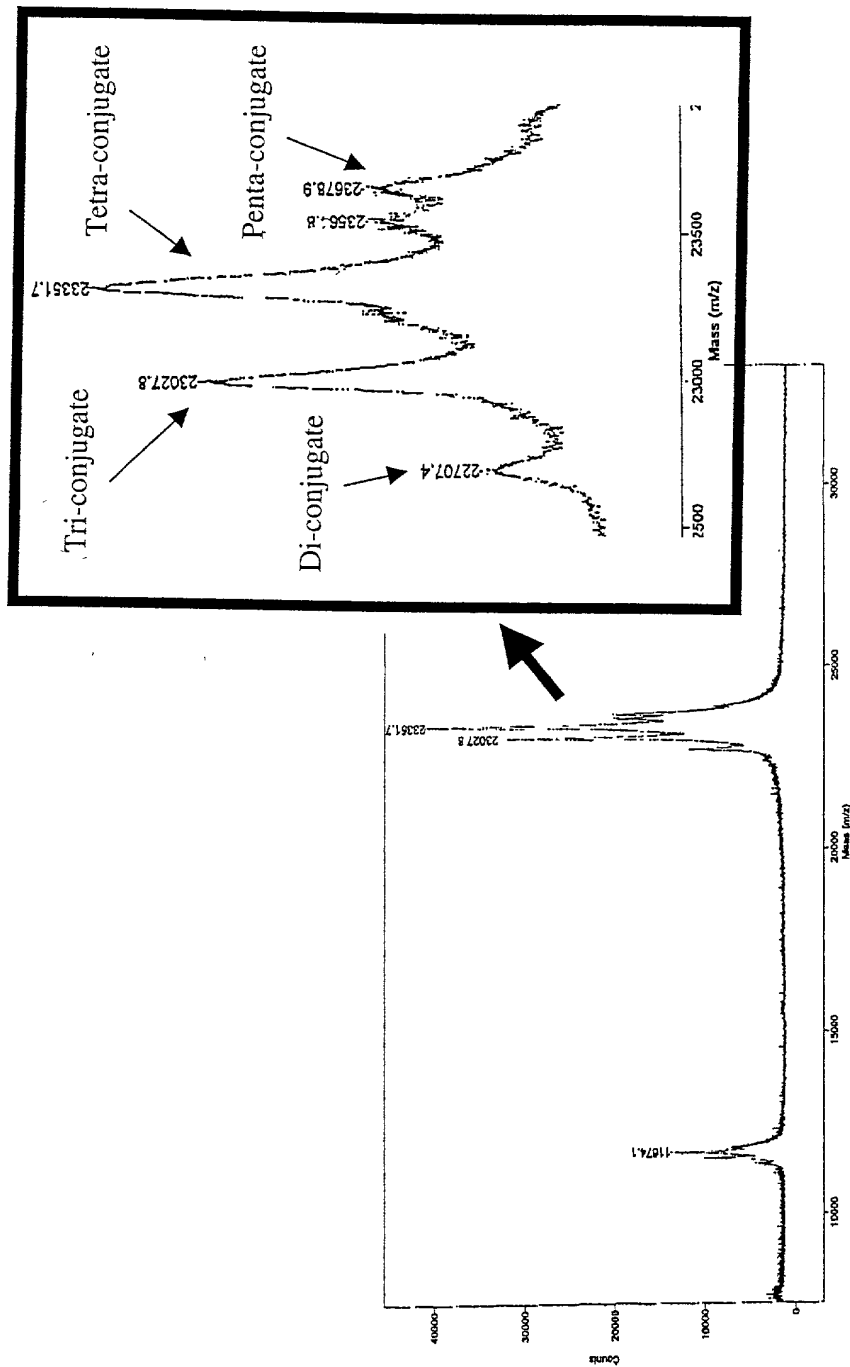


Figure 19

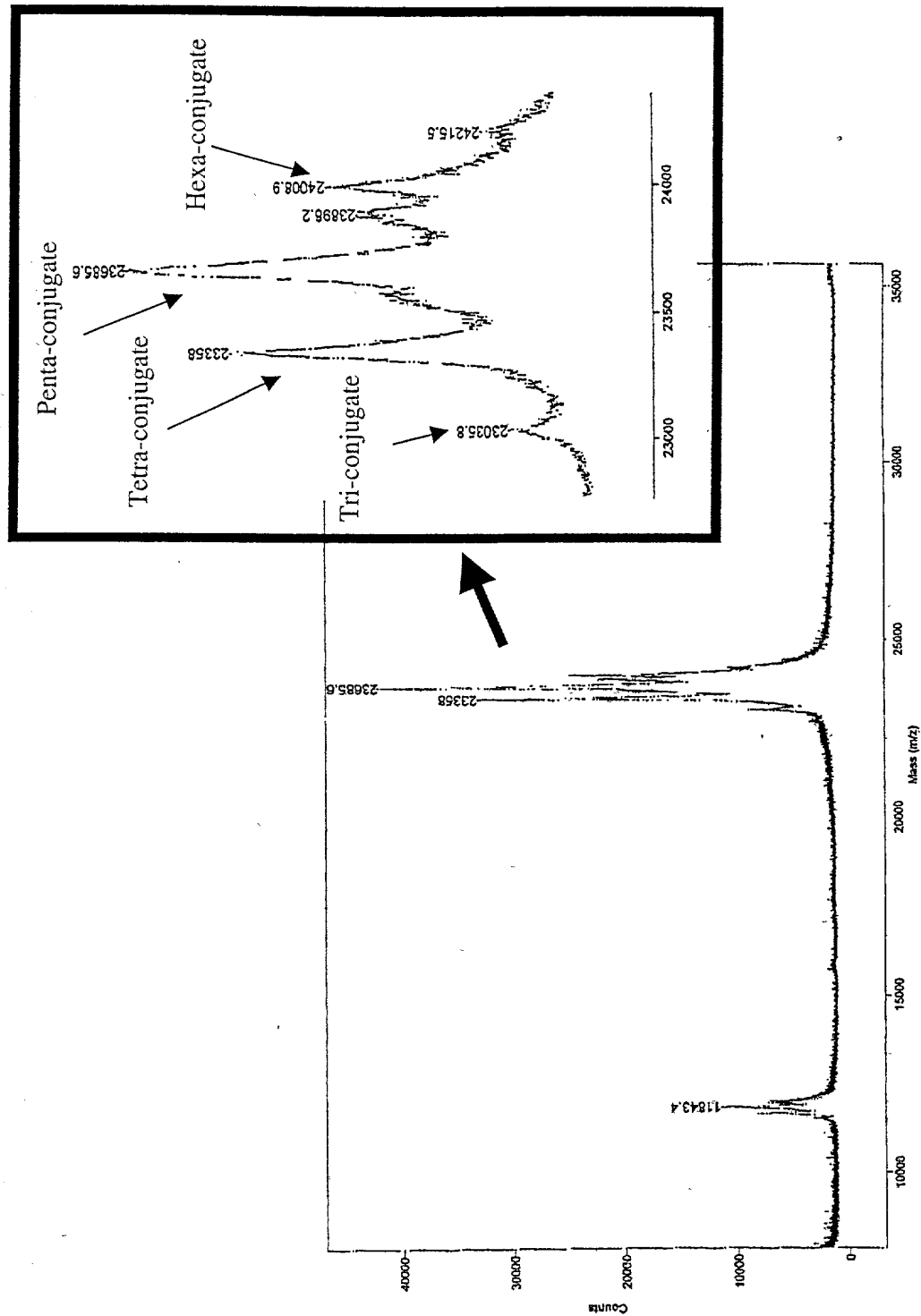


Figure 20

Mass spectrum of the hexa-conjugate + 8H⁺ complex. The x-axis is labeled 'm/z' and ranges from 2000 to 2450. The y-axis is labeled 'A' and ranges from 0 to 100. The base peak is at m/z 2406.5. Other significant peaks are labeled with their m/z values: 2396.0, 2400.0, 2407.8, 24113.4, 24173.5, 24180.8, 24200.8, 24224.1, 24232.3, 24250.4, and 24303.3. An arrow points to the peak at m/z 2406.5 with the label 'Hexa-conjugate + 8H⁺'.

Figure 21

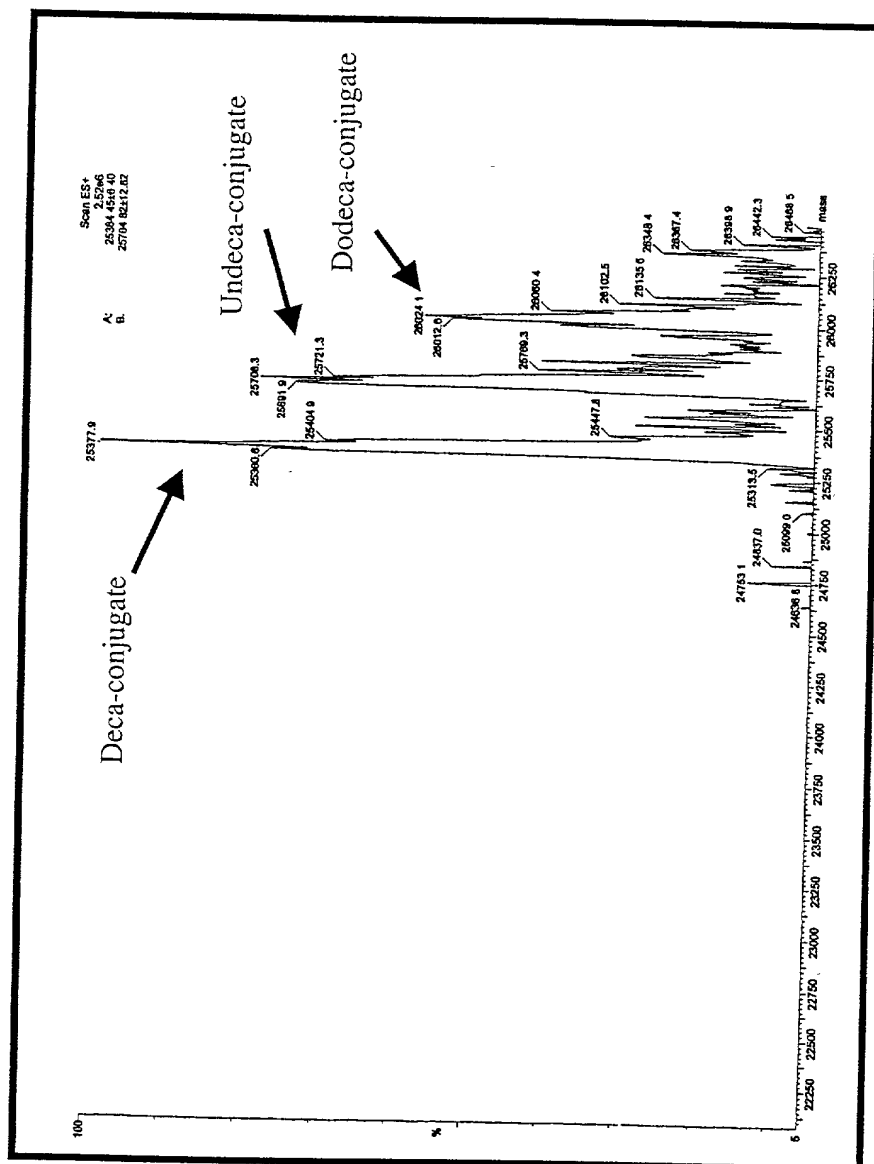


Figure 22

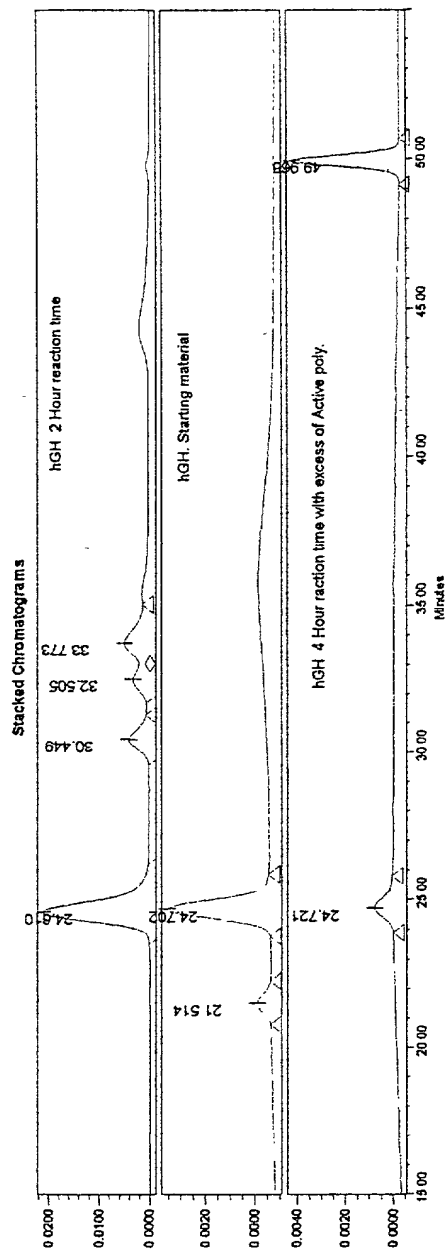


Figure 23

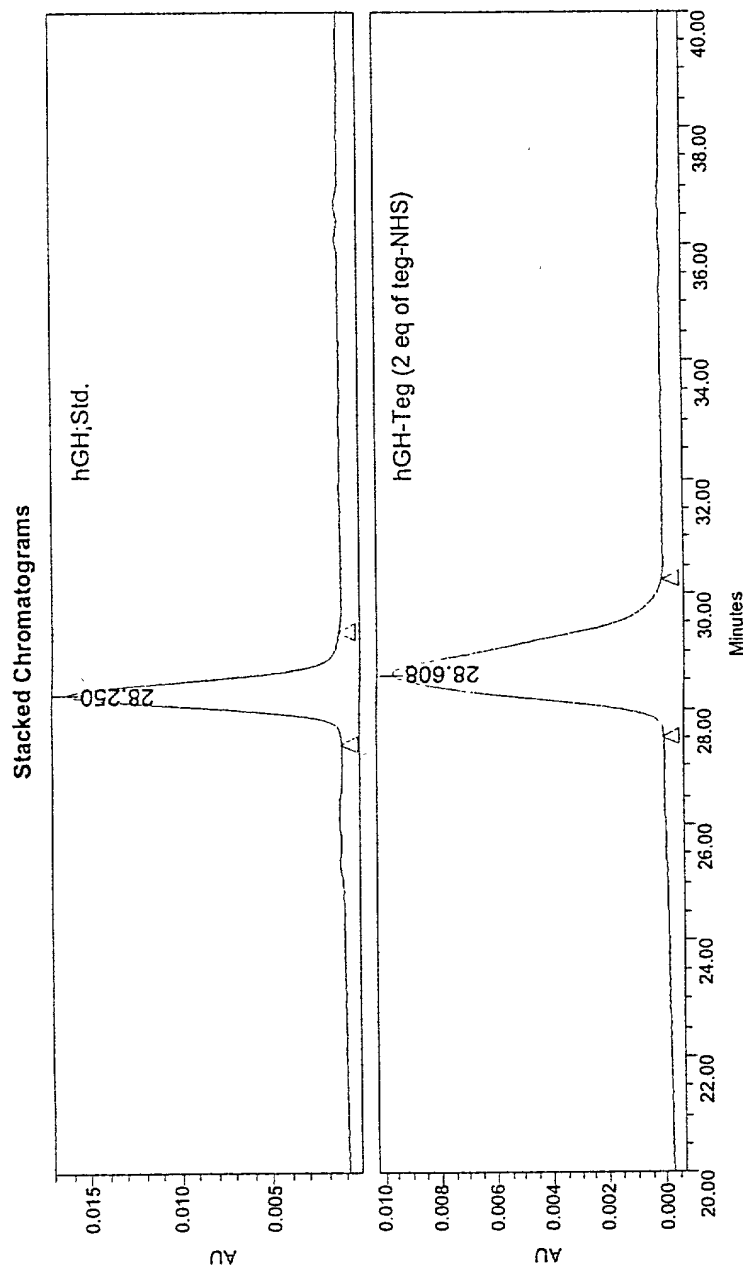


Figure 24

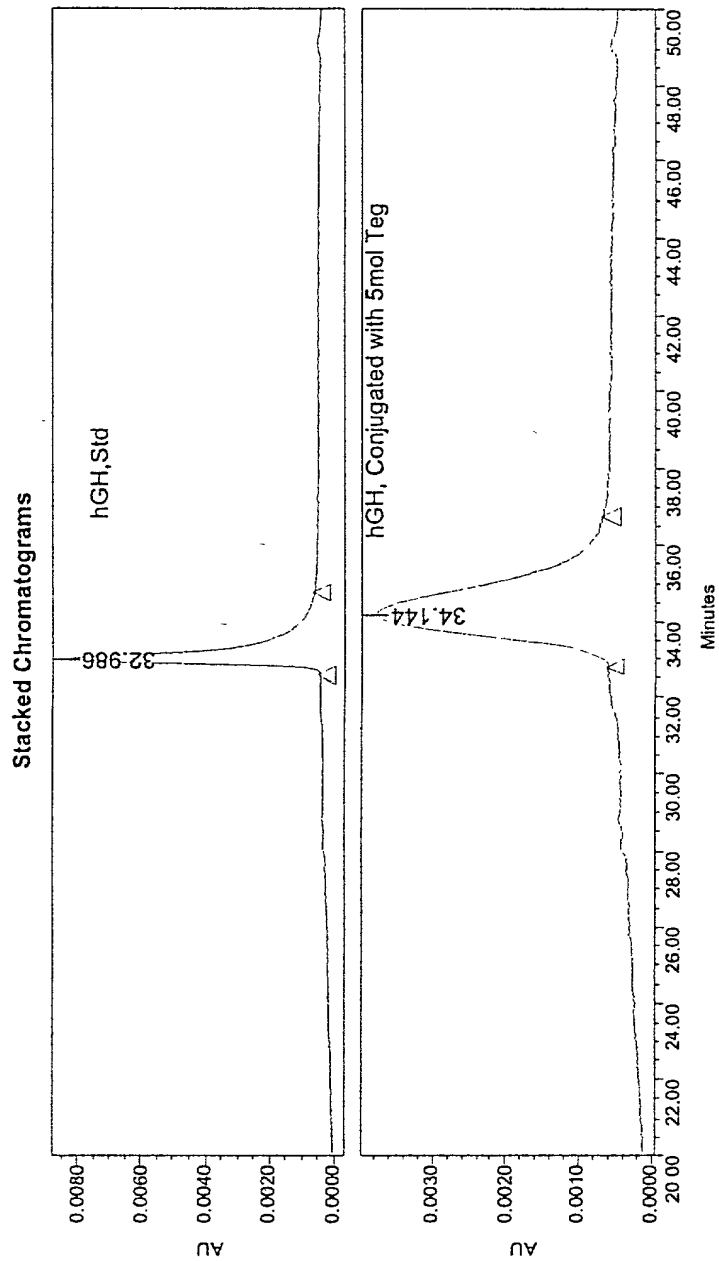


Figure 25

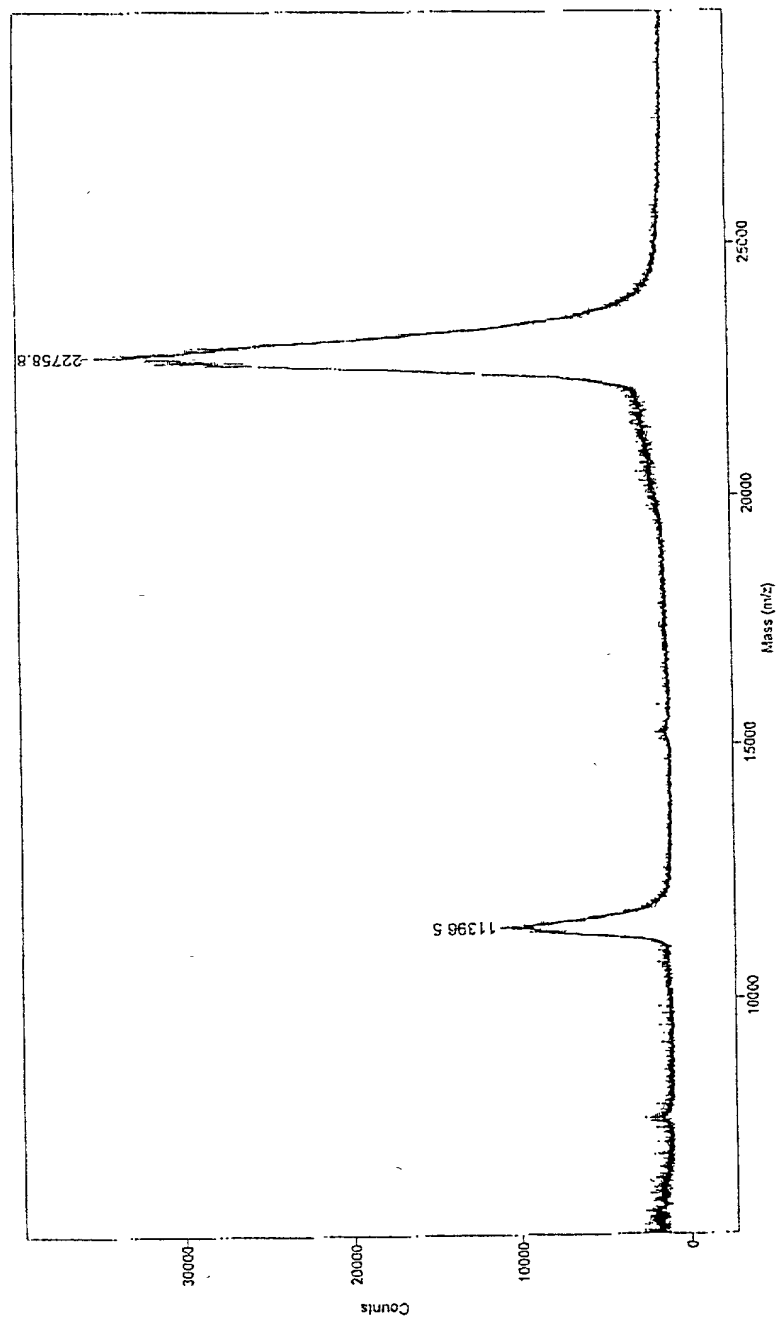


Figure 26

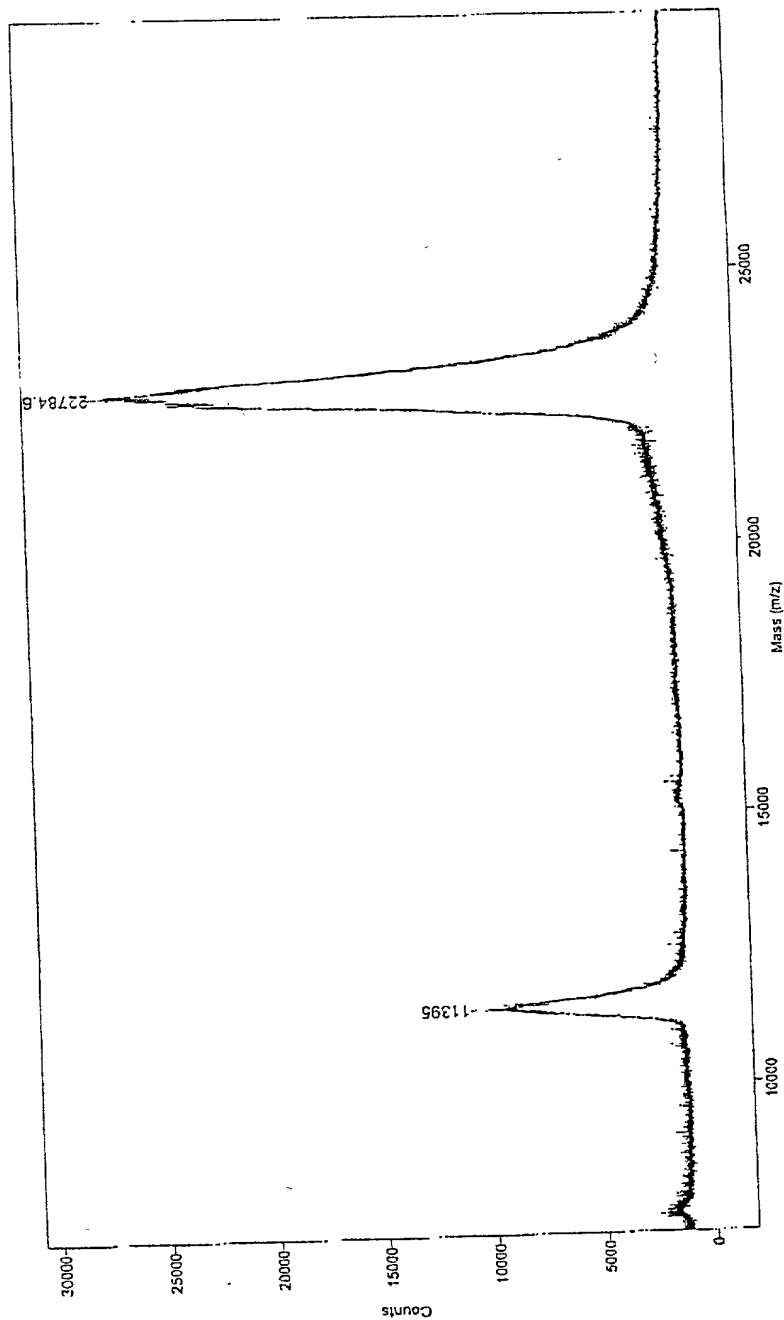


Figure 27

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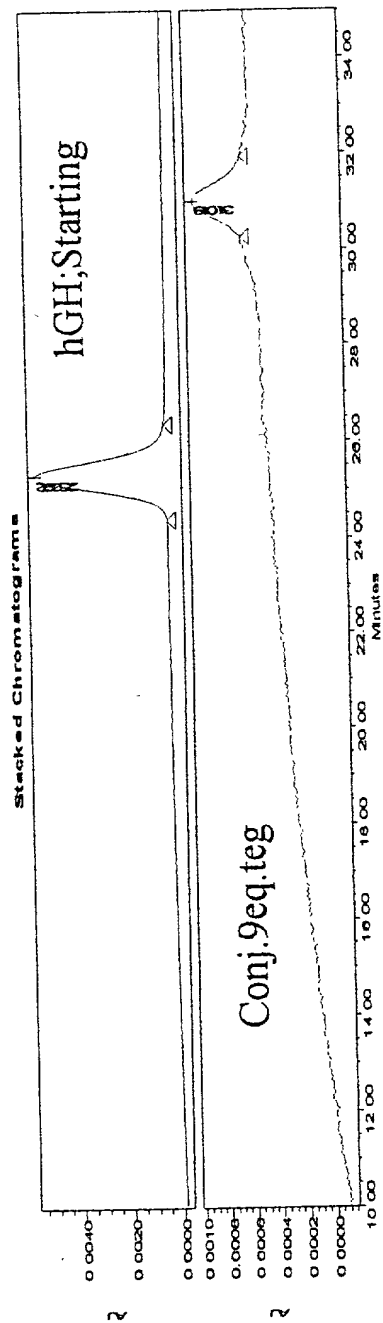


Figure 28

Figure 29

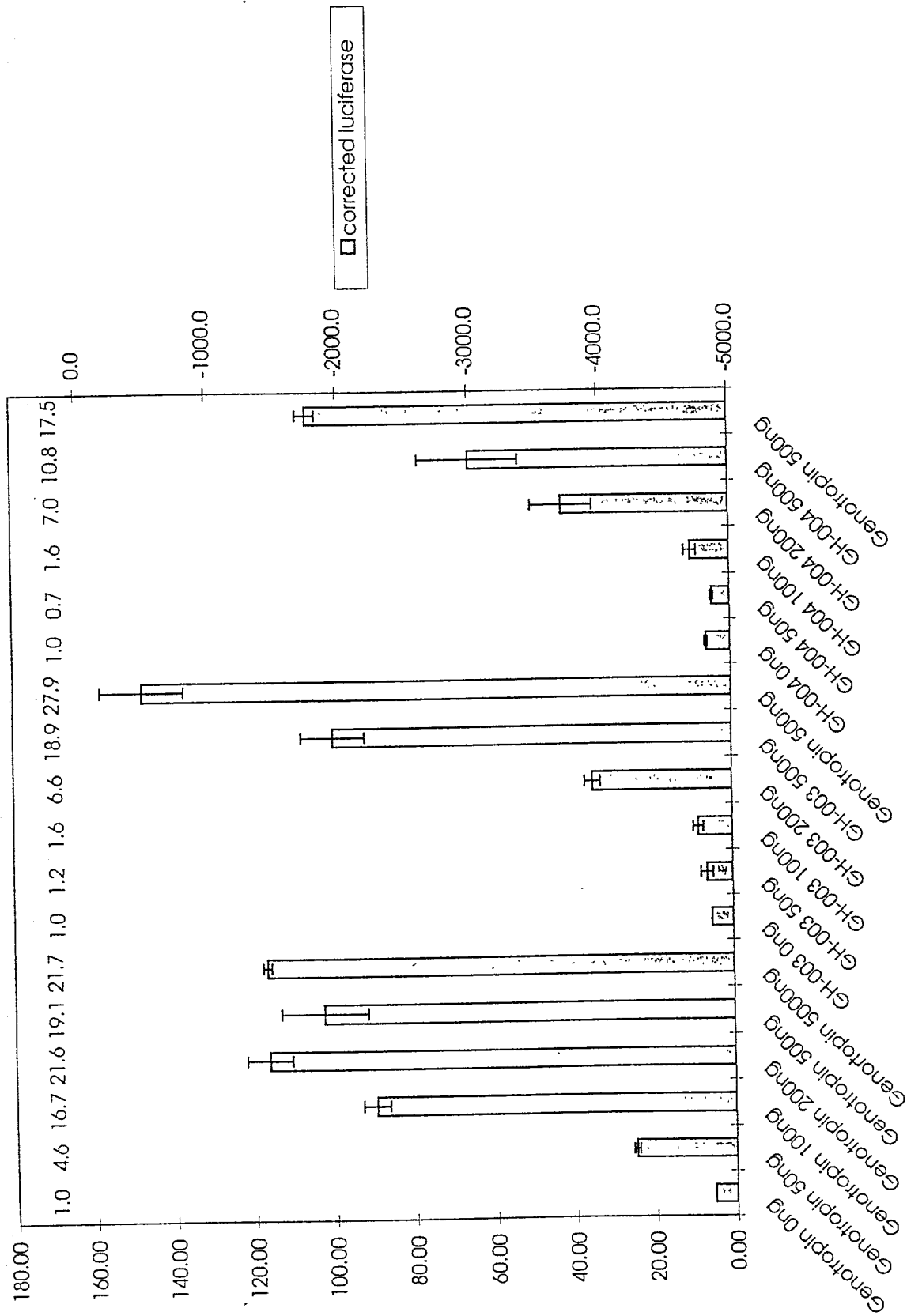


Figure 30